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0411

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O I P E

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/852,416

DATE: 04/16/2002
TIME: 16:00:02

Input Set : N:\CrF3\RULE60\09852416.raw
Output Set: N:\CRF3\04162002\I852416.raw

1 <110> APPLICANT: PKhosla, Chaitan
2 Ashley, Gary
3 Fu, Hong
4 Kao, Camilla M.
5 McDaniel, Robert
6 <120> TITLE OF INVENTION: COMBINATORIAL POLYKETIDE LIBRARIES
7 PRODUCED USING A MODULAR PKS GENE CLUSTER AS SCAFFOLD
8 <130> FILE REFERENCE: 30062-20005.02
9 <140> CURRENT APPLICATION NUMBER: 09/852,416
10 <141> CURRENT FILING DATE: 2001-05-09
12 <150> PRIOR APPLICATION NUMBER: 09/859,854
13 <151> PRIOR FILING DATE: 2001-05-16
16 <150> PRIOR APPLICATION NUMBER: PCT/US98/08792
17 <151> PRIOR FILING DATE: 1998-04-30
18 <150> PRIOR APPLICATION NUMBER: 60/076,919
19 <151> PRIOR FILING DATE: 1998-03-05
20 <150> PRIOR APPLICATION NUMBER: 08/846,247
21 <151> PRIOR FILING DATE: 1997-04-30
22 <150> PRIOR APPLICATION NUMBER: 08/486,645
23 <151> PRIOR FILING DATE: 1995-06-07
24 <150> PRIOR APPLICATION NUMBER: 08/238,811
25 <151> PRIOR FILING DATE: 1994-05-06
26 <160> NUMBER OF SEQ ID NOS: 44
27 <170> SOFTWARE: FastSEQ for Windows Version 4.0
29 <210> SEQ ID NO: 1
30 <211> LENGTH: 24
31 <212> TYPE: DNA
32 <213> ORGANISM: Artificial Sequence
33 <220> FEATURE:
34 <223> OTHER INFORMATION: Module 1 - A BamHI site engineered for the 5'
35 boundary of the acyltransferase domain.
36 <400> SEQUENCE: 1
37 g c g c a g c a g g g a t c c g t c t t c g t c
39 <210> SEQ ID NO: 2
40 <211> LENGTH: 24
41 <212> TYPE: DNA
42 <213> ORGANISM: Artificial Sequence
43 <220> FEATURE:
44 <223> OTHER INFORMATION: Module 1 - A PstI site engineered for introduction
45 between the acyltransferase and reductive domains.
46 <400> SEQUENCE: 2
47 c g c g t c t g g c t g c a g c c g a a g c c g
49 <210> SEQ ID NO: 3

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Input Set : N:\Crf3\RULE60\09852416.raw
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50 <211> LENGTH: 24
51 <212> TYPE: DNA
52 <213> ORGANISM: Artificial Sequence
53 <220> FEATURE:
54 <223> OTHER INFORMATION: Module 1 - A XbaI site engineered for introduction
55 at the 3' end of the reductive domain.
56 <400> SEQUENCE: 3
57 ggcgggtga gatctaagcc ggcc 24
59 <210> SEQ ID NO: 4
60 <211> LENGTH: 24
61 <212> TYPE: DNA
62 <213> ORGANISM: Artificial Sequence
63 <220> FEATURE:
64 <223> OTHER INFORMATION: Module 2 - A BamHI site engineered for the 5'
65 boundary of the acyltransferase domain.
66 <400> SEQUENCE: 4
67 tccgacggtg gatccgtgtt cgtc 24
69 <210> SEQ ID NO: 5
70 <211> LENGTH: 24
71 <212> TYPE: DNA
72 <213> ORGANISM: Artificial Sequence
73 <220> FEATURE:
74 <223> OTHER INFORMATION: Module 2 - A PstI site engineered for introduction
75 between the acyltransferase and reductive domains.
76 <400> SEQUENCE: 5
77 cggttctggc tgcagccgga ccgc 24
79 <210> SEQ ID NO: 6
80 <211> LENGTH: 24
81 <212> TYPE: DNA
82 <213> ORGANISM: Artificial Sequence
83 <220> FEATURE:
84 <223> OTHER INFORMATION: Module 2 - A XbaI site engineered for introduction
85 at the 3' end of the reductive domain.
86 <400> SEQUENCE: 6
87 gtcggccaga gatctcgaga ggca 24
89 <210> SEQ ID NO: 7
90 <211> LENGTH: 24
91 <212> TYPE: DNA
92 <213> ORGANISM: Artificial Sequence
93 <220> FEATURE:
94 <223> OTHER INFORMATION: Module 3 - A BamHI site engineered for the 5'
95 boundary of the acyltransferase domain.
96 <400> SEQUENCE: 7
97 gacgggcgcg gatccgtctt cctg 24
99 <210> SEQ ID NO: 8
100 <211> LENGTH: 24
101 <212> TYPE: DNA
102 <213> ORGANISM: Artificial Sequence
103 <220> FEATURE:

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104 <223> OTHER INFORMATION: Module 3 - A PstI site engineered for introduction
105      between the acyltransferase and reductive domains.
106 <400> SEQUENCE: 8
107      cgctactggc tgcagccgc cgca
108 <210> SEQ ID NO: 9
109 <211> LENGTH: 24
110 <212> TYPE: DNA
111 <213> ORGANISM: Artificial Sequence
112 <220> FEATURE:
113 <223> OTHER INFORMATION: Module 3 - A XbaI site engineered for introduction
114      at the 3' end of the reductive domain.
115 <400> SEQUENCE: 9
116      ctcggcaaca gatctgagcg gccca
117 <210> SEQ ID NO: 10
118 <211> LENGTH: 24
119 <212> TYPE: DNA
120 <213> ORGANISM: Artificial Sequence
121 <220> FEATURE:
122 <223> OTHER INFORMATION: Module 4 - A BamHI site engineered for the 5'
123      boundary of the acyltransferase domain.
124 <400> SEQUENCE: 10
125      gcgccgcgcg gatccgtcct ggtc
126 <210> SEQ ID NO: 11
127 <211> LENGTH: 24
128 <212> TYPE: DNA
129 <213> ORGANISM: Artificial Sequence
130 <220> FEATURE:
131 <223> OTHER INFORMATION: Module 4 - A PstI site engineered for introduction
132      between the acyltransferase and reductive domains.
133 <400> SEQUENCE: 11
134      cgcttctggc tgcagccgca ccgg
135 <210> SEQ ID NO: 12
136 <211> LENGTH: 24
137 <212> TYPE: DNA
138 <213> ORGANISM: Artificial Sequence
139 <220> FEATURE:
140 <223> OTHER INFORMATION: Module 4 - A XbaI site engineered for introduction
141      at the 3' end of the reductive domain.
142 <400> SEQUENCE: 12
143      ctcggccaga gatctaaggc cggg
144 <210> SEQ ID NO: 13
145 <211> LENGTH: 24
146 <212> TYPE: DNA
147 <213> ORGANISM: Artificial Sequence
148 <220> FEATURE:
149 <223> OTHER INFORMATION: Module 5 - A BamHI site engineered for the 5'
150      boundary of the acyltransferase domain.
151 <400> SEQUENCE: 13
152      actcgccgcg gatccgcgtt ggtg
  
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Input Set : N:\Crf3\RULE60\09852416.raw
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159 <210> SEQ ID NO: 14
160 <211> LENGTH: 24
161 <212> TYPE: DNA
162 <213> ORGANISM: Artificial Sequence
163 <220> FEATURE:
164 <223> OTHER INFORMATION: Module 5 - A PstI site engineered for introduction
165 between the acyltransferase and reductive domains.
166 <400> SEQUENCE: 14
167 cggtactggc tgcagatccc cacc
168 <210> SEQ ID NO: 15
169 <211> LENGTH: 24
170 <212> TYPE: DNA
171 <213> ORGANISM: Artificial Sequence
172 <220> FEATURE:
173 <223> OTHER INFORMATION: Module 5 - A XbaI site engineered for introduction
174 at the 3' end of the reductive domain.
175 <400> SEQUENCE: 15
176 gaccggctca gatctcgaaa ggag
177 <210> SEQ ID NO: 16
178 <211> LENGTH: 24
179 <212> TYPE: DNA
180 <213> ORGANISM: Artificial Sequence
181 <220> FEATURE:
182 <223> OTHER INFORMATION: Module 6 - A BamHI site engineered for the 5'
183 boundary of the acyltransferase domain.
184 <400> SEQUENCE: 16
185 tccggccggcg gatccgtttt cgtc
186 <210> SEQ ID NO: 17
187 <211> LENGTH: 24
188 <212> TYPE: DNA
189 <213> ORGANISM: Artificial Sequence
190 <220> FEATURE:
191 <223> OTHER INFORMATION: Module 6 - A PstI site engineered for introduction
192 between the acyltransferase and reductive domains.
193 <400> SEQUENCE: 17
194 cggtactggc tgcagccgga ggtg
195 <210> SEQ ID NO: 18
196 <211> LENGTH: 24
197 <212> TYPE: DNA
198 <213> ORGANISM: Artificial Sequence
199 <220> FEATURE:
200 <223> OTHER INFORMATION: Module 6 - A XbaI site engineered for introduction
201 at the 3' end of the reductive domain.
202 <400> SEQUENCE: 18
203 gacgtggcga gatctccggg ggtg
204 <210> SEQ ID NO: 19
205 <211> LENGTH: 23
206 <212> TYPE: DNA
207 <213> ORGANISM: Artificial Sequence
  
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PATENT APPLICATION: US/09/852,416 **TIME:** 16:00:02

Input Set : N:\CrF3\RULE60\09852416.raw
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213 <220> FEATURE:
214 <223> OTHER INFORMATION: A PstI site that is in-frame and upstream of XbaI
215 in pUC19 that generates this junction at the 5'
216 end of the cassette.
217 <400> SEQUENCE: 19
218 ctgcaggctcg actcttagcct ggt
220 <210> SEQ ID NO: 20
221 <211> LENGTH: 27
222 <212> TYPE: DNA
223 <213> ORGANISM: Artificial Sequence
224 <220> FEATURE:
225 <223> OTHER INFORMATION: Module rapAT2 (forward) Primer pairs used for PCR
226 amplification of rapamycin PKS cassettes.
227 <400> SEQUENCE: 20
228 tttagatctg tgttcgtctt cccgggt
230 <210> SEQ ID NO: 21
231 <211> LENGTH: 36
232 <212> TYPE: DNA
233 <213> ORGANISM: Artificial Sequence
234 <220> FEATURE:
235 <223> OTHER INFORMATION: Module rapAT2 (reverse) Primer pairs used for PCR
236 amplification of rapamycin PKS cassettes.
237 <400> SEQUENCE: 21
238 tttctgcagc cagtaccgct ggtgctggaa ggcgtta
240 <210> SEQ ID NO: 22
241 <211> LENGTH: 33
242 <212> TYPE: DNA
243 <213> ORGANISM: Artificial Sequence
244 <220> FEATURE:
245 <223> OTHER INFORMATION: Module rapAT14 (forward) Primer pairs used for PCR
246 amplification of rapamycin PKS cassettes.
247 <400> SEQUENCE: 22
248 ttggatccg ctttcctgtt cgacggcaa ggc
250 <210> SEQ ID NO: 23
251 <211> LENGTH: 33
252 <212> TYPE: DNA
253 <213> ORGANISM: Artificial Sequence
254 <220> FEATURE:
255 <223> OTHER INFORMATION: Module rapAT14 (reverse) Primer pairs used for PCR
256 amplification of rapamycin PKS cassettes.
257 <400> SEQUENCE: 23
258 tttctgcagc cagtaggact ggtgctggaa cgg
260 <210> SEQ ID NO: 24
261 <211> LENGTH: 36
262 <212> TYPE: DNA
263 <213> ORGANISM: Artificial Sequence
264 <220> FEATURE:
265 <223> OTHER INFORMATION: Module rapKR2 (forward) Primer pairs used for PCR
266 amplification of rapamycin PKS cassettes.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/852,416

DATE: 04/16/2002

TIME: 16:00:03

Input Set : N:\Crf3\RULE60\09852416.raw

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